
Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2008; month=4; day=22; hr=14; min=30; sec=32; ms=192;]

Validated By CRFValidator v 1.0.3

Application No: 10561671 Version No: 1.0

Input Set:

Output Set:

Started: 2008-04-08 18:52:13.349

Finished: 2008-04-08 18:52:15.498

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 149 ms

Total Warnings: 16

Total Errors: 0

No. of SeqIDs Defined: 22

Actual SeqID Count: 22

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Thr Trp Thr Val Thr Ile Pro Leu Leu Gln Gly Gln Asn Ile Gln Phe 50 55 60

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85 90 95

100 105 110

Gly Thr Glu Leu Val Asp Ala Val Glu Val Asn Pro Ser Asp Arg Asn 115 120 Gln Glu Ile Ser Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe Asp 135 Phe Pro Gly Arg Gly Asn Thr Tyr Ser Ser Phe Lys Trp Arg Trp Tyr 150 155 160 145 His Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn Arg Ile 170 175 165 Tyr Lys Phe Arg Gly Thr Gly Lys Ala Trp Asp Trp Glu Val Asp Thr 180 185 190 Glu Asn Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp 195 200 His Pro Glu Val Val Ser Glu Leu Lys Asn Trp Gly Lys Trp Tyr Val 210 215 220 Thr Thr Asn Ile Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile 235 225 230 Lys Tyr Ser Phe Phe Pro Asp Trp Leu Ser Tyr Val Arg Thr Gln Thr 250 245 255 Gln Lys Pro Leu Phe Ala Val Gly Glu Phe Trp Ser Tyr Asp Ile Ser 260 265 270 Lys Leu His Asn Tyr Ile Thr Lys Thr Asn Gly Ser Met Ser Leu Phe 280 275 Asp Ala Pro Leu His Asn Asn Phe Tyr Ile Ala Ser Lys Ser Gly Gly 290 295 300 Tyr Phe Asp Met Arg Thr Leu Leu Asn Asn Thr Leu Met Lys Asp Gln 315 305 310 Pro Thr Leu Ala Val Thr Leu Val Asp Asn His Asp Thr Glu Pro Gly

330

335

325

Gln Ser Leu Gln Ser Trp Val Glu Pro Trp Phe Lys Pro Leu Ala Tyr 345 Ala Phe Ile Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr Gly 355 360 365 Asp Tyr Tyr Gly Ile Pro Lys Tyr Asn Ile Pro Ala Leu Lys Ser Lys 370 375 380 Leu Asp Pro Leu Ile Ala Arg Asp Tyr Ala Tyr Gly Thr Gln 385 390 395 400 His Asp Tyr Ile Asp Ser Ala Asp Ile Ile Gly Trp Thr Arg Glu Gly 405 410 415 Val Ala Glu Lys Ala Asn Ser Gly Leu Ala Ala Leu Ile Thr Asp Gly 420 425 Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Gln His Ala Gly Lys 435 440 445 Thr Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn 460 455 450 Ala Asp Gly Trp Gly Glu Phe Lys Val Asn Gly Gly Ser Val Ser Ile 465 470 475 Trp Val Pro Lys <210> 5 <211> 485 <212> PRT <213> Bacillus sp <400> 5 Ala Asn Thr Ala Pro Val Asn Gly Thr Met Met Gln Tyr Phe Glu Trp

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Tyr Lys 50	Gly Thr	Ser Gln	Ala Asp 55	o Val	Gly	Tyr	Gly 60	Val	Tyr	Asp	Leu
Tyr Asp 65) Leu Gly	Glu Phe	Asn Glr	n Lys	Gly	Thr 75	Ile	Arg	Thr	Lys	Tyr 80
Gly Thr	Lys Thr	Gln Tyr 85	Leu Glr	n Ala	Ile 90	Gln	Ala	Ala	Lys	Ser 95	Ala
Gly Met	Gln Val	Tyr Ala	Asp Val	Val 105	Phe	Asn	His	Lys	Ala 110	Gly	Ala
Asp Ser	Thr Glu	Trp Val	Asp Ala		Glu	Val	Asn	Pro 125	Ser	Asn	Arg
Asn Gln		Ser Gly	Thr Tyr	Gln	Ile	Gln	Ala 140	Trp	Thr	Lys	Phe
Asp Phe	Pro Gly	Arg Gly		Tyr	Ser	Ser 155	Phe	Lys	Trp	Arg	Trp 160
Tyr His	Phe Asp	Gly Thr	Asp Tr	Asp	Glu 170	Ser	Arg	Lys	Leu	Asn 175	Arg
Ile Tyr	Lys Phe	Arg Gly	Thr Gly	y Lys 185	Ala	Trp	Asp	Trp	Glu 190	Val	Asp
Thr Glu	Asn Gly 195	Asn Tyr	Asp Tyr		Met	Phe	Ala	Asp 205	Leu	Asp	Met
Asp His		Val Val	Ala Glu 215	ı Leu	Lys	Asn	Trp 220	Gly	Lys	Trp	Tyr
Val Asn 225	Thr Thr	Asn Val		y Phe	Arg	Leu 235	Asp	Ala	Val	Lys	His 240
Ile Lys	Tyr Ser	Phe Phe	Pro Asp	Trp	Leu 250	Ser	Tyr	Val	Arg	Asn 255	Gln

Thr Gly Lys Asn Leu Phe Ala Val Gly Glu Phe Trp Gly Tyr Asp Val

260 265 270

Asn Lys Leu His Asn Tyr Ile Thr Lys Thr Asn Gly Ala Met Ser Leu 275 280 285

Phe Asp Ala Pro Leu His Asn Asn Phe Tyr Ile Ala Ser Lys Ser Ser 290 295 300

Gly Tyr Phe Asp Met Arg Tyr Leu Leu Asn Asn Thr Leu Met Lys Asp 305 310 315

Gln Pro Ala Leu Ala Val Thr Leu Val Asp Asn His Asp Thr Gln Pro 325 330 335

Gly Gln Ser Leu Gln Ser Trp Val Glu Pro Trp Phe Lys Pro Leu Ala 340 345 350

Tyr Ala Phe Ile Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr 355 360 365

Gly Asp Tyr Tyr Gly Ile Pro Lys Tyr Asn Ile Pro Gly Leu Lys Ser 370 375 380

Gln Arg Asp Tyr Ile Asp His Gln Asp Ile Ile Gly Trp Thr Arg Glu
405 410 415

Gly Ile Asp Ala Lys Pro Asn Ser Gly Leu Ala Ala Leu Ile Thr Asp 420 425 430

Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Arg His Ala Gly 435 440 445

Lys Val Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile 450 455 460

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Ser Leu Ala Asn Leu Gly Ile Thr Ala Leu Trp Leu Pro Pro Ala Tyr
             40
Lys Gly Thr Ser Ser Ser Asp Val Gly Tyr Gly Val Tyr Asp Leu Tyr
     55 60
  50
Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Val Arg Thr Lys Tyr Gly
         70
                           75
65
Thr Lys Thr Gln Tyr Ile Gln Ala Ile Gln Ala Ala His Thr Ala Gly
          85 90 95
Met Gln Val Tyr Ala Asp Val Val Phe Asn His Lys Ala Gly Ala Asp
      100 105 110
Gly Thr Glu Leu Val Asp Ala Val Glu Val Asn Pro Ser Asp Arg Asn
   115 120 125
Gln Glu Ile Ser Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe Asp
  130
        135
Phe Pro Gly Arg Gly Asn Thr Tyr Ser Ser Phe Lys Trp Arg Trp Tyr
                           155
145
            150
                                       160
His Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn Arg Ile
         165 170 175
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Tyr Lys Phe Arg Gly Thr Gly Lys Ala Trp Asp Trp Glu Val Asp Thr 185

180

Glu	Asn	Gly 195	Asn	Tyr	Asp	Tyr	Leu 200	Met	Tyr	Ala	Asp	Leu 205	Asp	Met	Asp
His	Pro 210	Glu	Val	Val	Ser	Glu 215	Leu	Lys	Asn	Trp	Gly 220	Lys	Trp	Tyr	Val
Ile 225	Thr	Thr	Asn	Ile	Asp 230	Gly	Phe	Arg	Leu	Asp 235	Ala	Val	Lys	His	Ile 240
Lys	Tyr	Ser	Phe	Phe 245	Pro	Asp	Trp	Leu	Ser 250	Tyr	Leu	Arg	Thr	Gln 255	Thr
			Leu 260					265					270		
		275	Asn				280					285			
	290					295					300				Gly
305			Met		310					315					320
			Gln	325					330					335	
			340 Leu					345					350		
		355	Gly				360					365			
	370		Leu			375					380				
385			Ile		390					395					400
****	1121	- y -	110	405	11011	1114	1127	110	410	CLY			*****	415	<u>y</u>

Val Ala Glu Lys Ala Asn Ser Gly Leu Ala Ala Leu Ile Thr Asp Gly

420 425 430

Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Gln His Ala Gly Lys
435 440 445

Thr Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn 450 450 460

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Trp Val Pro Lys

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Ala Gly Ser Val Pro Val Asn Gly Thr Met Met Gln Tyr Phe Glu Trp 35 40 45

Tyr Leu Pro Asp Asp Gly Thr Leu Trp Thr Lys Val Ala Asn Asn Ala 50 55 60

Gln Ser Leu Ala Asn Leu Gly Ile Thr Ala Leu Trp Leu Pro Pro Ala 65 70 75 80

Tyr Lys Gly Thr Ser Ser Ser Asp Val Gly Tyr Gly Val Tyr Asp Leu 85 90 95

Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Val Arg Thr Lys Tyr
100 105 110

Gly Thr Lys Thr Gln Tyr Ile Gln Ala Ile Gln Ala Ala His Thr Ala 115 120 125

Gly Met Gln Val Tyr Ala Asp Val Val Phe Asn His Lys Ala Gly Ala 130 $$135\$

Asn Gln Glu Ile Ser Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe 165 170 175

Asp Phe Pro Gly Arg Gly Asn Thr Tyr Ser Ser Phe Lys Trp Arg Trp 180 185 190

Tyr His Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn Arg 195 200 205

Ile Tyr Lys Phe Arg Gly Thr Gly Lys Ala Trp Asp Trp Glu Val Asp 210 215 220

Thr Glu Asn Gly Asn Tyr As